

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/672,177A
Source: TFNO
Date Processed by STIC: 2-4-05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
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Revised 01/24/05



IFWO

RAW SEQUENCE LISTING

DATE: 02/04/2005

PATENT APPLICATION: US/10/672,177A

TIME: 16:27:06

Input Set : N:\FATIMA\10672177.txt

Output Set: N:\CRF4\02042005\J672177A.raw

W--> 2 Alpha. ST25.txt

5 <110> APPLICANT: AlphaMed Pharmaceuticals Corporation

8 <120> TITLE OF INVENTION: DNA for Expression of Alpha I-Antitrypsin in Methylootropic Yeast

11 <130> FILE REFERENCE: 110113200

C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/672,177A

C--> 16 <141> CURRENT FILING DATE: 2003-09-26

19 <150> PRIOR APPLICATION NUMBER: PCT/US01/50677

21 <151> PRIOR FILING DATE: 2001-12-21

24 <160> NUMBER OF SEQ ID NOS: 8

27 <170> SOFTWARE: PatentIn version 3.1

Does Not Comply
Corrected Diskette Needed

ERRORED SEQUENCES

59 <210> SEQ ID NO: 3

61 <211> LENGTH: 24

63 <212> TYPE: DNA

65 <213> ORGANISM: Homo sapiens

68 <400> SEQUENCE: 3

E--> 69 cagaagacag atacatcca ccat, 24

VERIFICATION SUMMARY

DATE: 02/04/2005

PATENT APPLICATION: US/10/672,177A

TIME: 16:27:07

Input Set : N:\FATIMA\10672177.txt

Output Set: N:\CRF4\02042005\J672177A.raw

L:2 M:259 W: Allowed number of lines exceeded, (1) GENERAL INFORMATION:
L:14 M:270 C: Current Application Number differs, Replaced Current Application Number
L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:47 M:259 W: Allowed number of lines exceeded, <211> LENGTH:
L:69 M:254 E: No. of Bases conflict, LENGTH: Input:24 Counted:23 SEQ:3
L:69 M:252 E: No. of Seq. differs, <211> LENGTH: Input:24 Found:23 SEQ:3
L:94 M:259 W: Allowed number of lines exceeded, <213> ORGANISM: